

## RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/509,773

Source: PG/10

Date Processed by STIC: 7/7/05

# ***ENTERED***

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PCT

## RAW SEQUENCE LISTING

DATE: 07/07/2005

PATENT APPLICATION: US/10/509,773

TIME: 13:28:01

Input Set : D:\seqlist.txt

Output Set: N:\CRF4\07072005\J509773.raw

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3 <110> APPLICANT: Delaney, Allen
5 <120> TITLE OF INVENTION: Cancer Associated Protein Phosphatases and their
6   uses
8 <130> FILE REFERENCE: SMAR-044
10 <140> CURRENT APPLICATION NUMBER: 10/509,773
C--> 11 <141> CURRENT FILING DATE: 2004-09-28
13 <150> PRIOR APPLICATION NUMBER: CA03/00393
14 <151> PRIOR FILING DATE: 2003-03-19
16 <150> PRIOR APPLICATION NUMBER: 60/368,859
17 <151> PRIOR FILING DATE: 2002-03-28
19 <160> NUMBER OF SEQ ID NOS: 12
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30 <222> LOCATION: (0)...(0)
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36 gcgaggaggg acgctggggc tgcccgtgcg gcacgggggc ggggaccggc aaggcgggac      180
37 catttcccg cataggtctc ggtgcccctg cccggctccc gccgggaagt tctaggccgc      240
38 cgcacagaaa gccctgccct ccacgccggg tctctggagc gccctgggtt gcccggccgg      300
39 tccctgccgc tgacttggtg aactgcgag cactcagtc ctcgcgcgc ctcctcccc      360
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44 gtgcatccca gcagcggatt caccatctca aaacctgaca agacatttca aagaaagtat      660
45 taaattcatt cacgagtgcc ggctccgcgg tgagagctgc cttgtacact gcctggccgg      720
46 ggtctccagg agcgtgacac tggatgatcg atacatcatg accgtcactg actttggctg      780
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48 ccagagacag ctccaggagt ttgagaagca tgagggtccat cagtatcggc agtggctgaa      900
49 ggaagaatat ggagagagcc ctttgcagga tgcagaagaa gccaaaaaca ttctggccgc      960
50 tccaggaatt ctgaagttct gggcctttct cagaagactg taatgtacct gaagtttctg      1020
51 aaatattgca aaccacacaga gtttaggctg gtgctgccaa aaagaaaagc aacatagagt      1080
52 ttaagtatcc agtagtgatt tgtaaaactt tttttcattt gaagctgaat atatacgtag      1140
53 tcatgtttat gttgagaact aaggatatct tttagcaaga gaaaatattt tccccttatc      1200
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56 gacgagccca gtgtagtgt gtggcgtgaa ctctgcccgt gtgttctcaa attccccagc 1380
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62 &lt;210&gt; SEQ ID NO: 2

63 &lt;211&gt; LENGTH: 184

64 &lt;212&gt; TYPE: PRT

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68 &lt;221&gt; NAME/KEY: UNSURE

69 &lt;222&gt; LOCATION: (0)...(0)

70 &lt;223&gt; OTHER INFORMATION: MKPX polypeptide

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75 Phe Lys Asp Ala Arg Asp Ala Glu Gln Leu Ser Lys Asn Lys Val Thr
76 20 25 30
77 His Ile Leu Ser Val His Asp Ser Ala Arg Pro Met Leu Glu Gly Val
78 35 40 45
79 Lys Tyr Leu Cys Ile Pro Ala Ala Asp Ser Pro Ser Gln Asn Leu Thr
80 50 55 60
81 Arg His Phe Lys Glu Ser Ile Lys Phe Ile His Glu Cys Arg Leu Arg
82 65 70 75 80
83 Gly Glu Ser Cys Leu Val His Cys Leu Ala Gly Val Ser Arg Ser Val
84 85 90 95
85 Thr Leu Val Ile Ala Tyr Ile Met Thr Val Thr Asp Phe Gly Trp Glu
86 100 105 110
87 Asp Ala Leu His Thr Val Arg Ala Gly Arg Ser Cys Ala Asn Pro Asn
88 115 120 125
89 Val Gly Phe Gln Arg Gln Leu Glu Phe Glu Lys His Glu Val His
90 130 135 140
91 Gln Tyr Arg Gln Trp Leu Lys Glu Glu Tyr Gly Glu Ser Pro Leu Gln
92 145 150 155 160
93 Asp Ala Glu Glu Ala Lys Asn Ile Leu Ala Ala Pro Gly Ile Leu Lys
94 165 170 175
95 Phe Trp Ala Phe Leu Arg Arg Leu
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102 &lt;213&gt; ORGANISM: Homo sapiens

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105 &lt;221&gt; NAME/KEY: misc\_feature

106 &lt;222&gt; LOCATION: (0)...(0)

107 &lt;223&gt; OTHER INFORMATION: PTP4A1 polynucleotide

109 &lt;400&gt; SEQUENCE: 3

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111 tgctccgagc cgctcactgc atggtagagt ctggtgccc cgccgccgcc tgcacgccg 120
112 ccaccgccgc tccgccacga ccaccgccgc ctcttgtcc tgcagccacc gccaccgcct 180

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115 gattactgcc aggcacagca cgacctctat gcagacaagt gaactgtaga aactgattac 360
116 tgctccacca agaagccccc ataagagtgg ttatcctgga cacagaagtg ttgaaatcca 420
117 cagagcattt tacaagagtt ctgacctgga tggggtaaac ctgagtgcac ttcttttctg 480
118 ttggcctcag tattactgga ttgaagaatt gctgcttctt gttaggaggt tcatttcact 540
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120 tagacttcag tttctttgga tcatttctgt attcaatttt ttttaattatt tcataaccct 660
121 attgagtgtt ttttaactaa attaacatgg ctgcaatgaa ccgcccagct cctgtggaag 720
122 tcacatacaa gaacatgaga tttcttatta cacacaatcc aaccaatgag accttaaaca 780
123 aatttataga ggaacttaag aagtatggag ttaccacaat agtaagagta tgtgaagcaa 840
124 cttatgacac tactcttgtg gagaaagaag gtatccatgt tcttgattgg ccttttgatg 900
125 atggtgcacc accatccaac cagattgttg atgactgggt aagtcttgtg aaaattaaagt 960
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127 cagtacttgt tgccctagca ttaattgaag gtggaatgaa atacgaagat gcagtacaat 1080
128 tcataagaca aaagcggcgt ggagctttta acagcaagca acttctgtat ttggagaagt 1140
129 atcgctcctaa aatgcggctg cgtttcaaag attccaacgg tcatagaaac aactgttgca 1200
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155 agataatttt gaaccaattt attattgtgt actgaggaga aataatgtat agtagaggac 2760
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158 atattagtat ctggctacat gatggccatg tcaagt 2916
161 <210> SEQ ID NO: 4
162 <211> LENGTH: 173
163 <212> TYPE: PRT

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164 <213> ORGANISM: Homo sapiens  
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 167 <221> NAME/KEY: UNSURE  
 168 <222> LOCATION: (0)...(0)  
 169 <223> OTHER INFORMATION: PTP4A1 polypeptide sequence  
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 175 20 25 30  
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 177 35 40 45  
 178 Cys Glu Ala Thr Tyr Asp Thr Thr Leu Val Glu Lys Glu Gly Ile His  
 179 50 55 60  
 180 Val Leu Asp Trp Pro Phe Asp Asp Gly Ala Pro Pro Ser Asn Gln Ile  
 181 65 70 75 80  
 182 Val Asp Asp Trp Leu Ser Leu Val Lys Ile Lys Phe Arg Glu Glu Pro  
 183 85 90 95  
 184 Gly Cys Cys Ile Ala Val His Cys Val Ala Gly Leu Gly Arg Ala Pro  
 185 100 105 110  
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 187 115 120 125  
 188 Ala Val Gln Phe Ile Arg Gln Lys Arg Arg Gly Ala Phe Asn Ser Lys  
 189 130 135 140  
 190 Gln Leu Leu Tyr Leu Glu Lys Tyr Arg Pro Lys Met Arg Leu Arg Phe  
 191 145 150 155 160  
 192 Lys Asp Ser Asn Gly His Arg Asn Asn Cys Cys Ile Gln  
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196 &lt;210&gt; SEQ ID NO: 5

197 &lt;211&gt; LENGTH: 2759

198 &lt;212&gt; TYPE: DNA

199 &lt;213&gt; ORGANISM: Homo sapiens

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202 &lt;221&gt; NAME/KEY: misc\_feature

203 &lt;222&gt; LOCATION: (0)...(0)

204 &lt;223&gt; OTHER INFORMATION: PTPN7 polynucleotide sequence

206 &lt;400&gt; SEQUENCE: 5

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 213 agccacccag ccccaagcaa ctggaagaag aattcttgaa gatcccttca aactttgtca 420  
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 215 caaatcccca gagccgtgtc tgtctaggcc gggcacagag ccaggaggac ggagattaca 540  
 216 tcaatgccaa ctacatccga ggctatgacg ggaaggagaa ggtctacatt gccaccagc 600  
 217 gcccctatgcc caacactgtg tcggacttct gggagatggg gtggcaagag gaagtgtccc 660  
 218 tcattgtcat gctcactcag ctccgagagg gcaaggagaa atgtgtccac tactggccca 720

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221 tcctcttttc ggcttgcca gaccatcaga caccagaatc agctgggccc ctgctgcgcc 900
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225 gaggggggat gatccagacg gcagagcagt accagttcct gcaccacact ttggccctgt 1140
226 atgcaggcca gctgcctgag gaacccagcc cctgaccctc gccaccctcc ggtggcccag 1200
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255 &lt;210&gt; SEQ ID NO: 6

256 &lt;211&gt; LENGTH: 339

257 &lt;212&gt; TYPE: PRT

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260 &lt;220&gt; FEATURE:

261 &lt;221&gt; NAME/KEY: UNSURE

262 &lt;222&gt; LOCATION: (0)...(0)

263 &lt;223&gt; OTHER INFORMATION: PTPN7 polypeptide sequence

265 &lt;400&gt; SEQUENCE: 6

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268 Leu Gln Glu Arg Gly Ser Asn Val Ala Leu Met Leu Asp Val Arg
269 20 25 30
270 Ser Leu Gly Ala Val Glu Pro Ile Cys Ser Val Asn Thr Pro Arg Glu
271 35 40 45

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L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date